

## STIC-Biotech/ChemLib

114121

**From:** Epps-Ford, Janet  
**Sent:** Wednesday, February 11, 2004 11:14 AM  
**To:** STIC-Biotech/ChemLib  
**Subject:** FW: Amino acid sequence search

Application: 09/914,020  
SEQ ID NO: 3

Type of search: Oligo search of SEQ ID NO: 3 in pending and issued amino acid databases.

Thanks.

-----Original Message-----

**From:** Schreiber, David  
**Sent:** Wednesday, February 11, 2004 11:05 AM  
**To:** Epps-Ford, Janet  
**Subject:** RE: Amino acid sequence search

Just request an oligo search of seq 3. Do not specify a word size since 10 is already a relatively low probability event statistically.

-----Original Message-----

**From:** Epps-Ford, Janet  
**Sent:** Wednesday, February 11, 2004 11:00 AM  
**To:** Schreiber, David  
**Subject:** Amino acid sequence search

Hi David, I am examining the following claim:

A method according to claim 1 wherein said compound is a peptide having a sequence comprising from about 10 to about 50 contiguous amino acids from SEQ ID NO: 3.

Application: 09/914,020.

I am not sure how to request the appropriate search, what would you recommend??

*Thanks,*

*Janet L. Epps-Ford, Ph.D.  
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✓ Pending Nucleic Acid and Pending Amino Acid database searches generate two sets of results each. The Pending databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches. Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions .rnpm and .rnpn. Searches run against the Amino Acid Pending database produce two sets of results, with the extensions .rapm and .rapn.

***Because they contain data that is confidential, the results of Pending database searches should not be left in the case .***